

AMENDMENTS TO THE SPECIFICATION

Please make the following amendments to the specification.

Please replace paragraph 0146 with the following paragraph:

Reference is now made to Fig. 12A which is an annotated sequence of an EST comprising a novel gene detected by the gene detection system of the present invention. Fig. 12A shows the nucleotide sequence of a known human non-protein coding EST (Expressed Sequence Tag), identified as EST72223 (SEQ_ID_NO: 46756). It is appreciated that the sequence of this EST comprises sequences of one known miRNA gene, identified as MIR98, and of one novel GAM gene, referred to here as GAM24, detected by the bioinformatic gene detection system of the present invention, described hereinabove with reference to Fig. 2.

Please replace paragraph 0151 with the following paragraph:

Transcript preparations: Digoxigenin (DIG) labeled transcripts were prepared from EST72223 (TIGER), MIR98 and predicted precursor hairpins by using a DIG RNA labeling kit (Roche Molecular Biochemicals) according to the manufacturer's protocol. Briefly, PCR products with T7 promoter at the 5" end or T3 promoter at the 3" end were prepared from each DNA in order to use it as a template to prepare sense and antisense transcripts, respectively. MIR-98 was amplified using EST72223 as a templet template with T7miR98 forward primer: 5"-TAATACGACTCACTATAGGGTGGAGGTAGTAAGTTGTA TTGTT-3" (SEQ_ID_NO: 46760) and T3miR98 esse reverse primer: 5"-AATTAACCCCTCACTAAAGGGAAAGTAGTAGTAAG TTGTATAGTT-3" (SEQ_ID_NO: 46761). EST72223 was amplified with T7-EST 72223 forward primer: 5"-TAATACGACTCACTATAGGCCCTATTAGAGGGATTCTGCT -3" (SEQ_ID_NO: 46762) and T3-EST72223 reverse primer: 5"-AATTAACCCCTCACTAAAGGTTTTTTCTGAGACAG AGT3" (SEQ_ID_NO: 46763). Bet-4 was amplified using EST72223 as a templet template with Bet-4 forward primer: 5"-GAGGCAGGAGAAATGCTGA-3" (SEQ_ID_NO: 46764) and T3-EST72223 reverse primer: 5"-AATTAACCCCTCACTAA AGGCCCTGAGACAGAGTCITGCTC-3" (SEQ_ID_NO: 46765). The PCR products were cleaned and used for DIG-labeled or unlabeled transcription reactions with the appropriate polymerase. For transfection experiments, CAP reaction was performed by using a mMassage mMessage mMachine kit (Ambion).

Please replace paragraph 0157 with the following paragraph:

Reference is now made to Fig. 13A which is an annotated sequence of an EST comprising a novel gene detected by the gene detection system of the present invention. Fig. 13A shows the nucleotide sequence of a known human non-protein coding EST (Expressed Sequence Tag), identified as EST 7929020 (SEQ ID NO: 46757). It is appreciated that the sequence of this EST comprises sequences of two novel GAM genes, referred to here as GAM23 and GAM25, detected by the bioinformatic gene detection system of the present invention, described hereinabove with reference to Fig. 2.

Please replace paragraph 0160 with the following paragraph:

Reference is now made to Fig. 14A which is an annotated sequence of an EST comprising a novel gene detected by the gene detection system of the present invention. Fig. 14A shows the nucleotide sequence of a known human non-protein coding EST (Expressed Sequence Tag), identified as EST 1388749 (SEQ ID NO: 46758). It is appreciated that the sequence of this EST comprises sequence of a novel GAM gene, referred to here as GAM26, detected by the bioinformatic gene detection system of the present invention, described hereinabove with reference to Fig. 2.